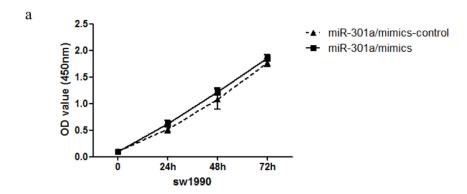
MicroRNA-301a-3p promotes pancreatic cancer progression via negative regulation of *SMAD4*

Supplementary Material



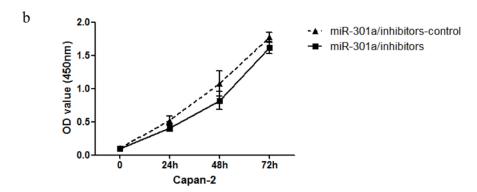


Figure S1: Role of miR-301a-3p in proliferation. Cell proliferation was measured by CCK-8 assay in sw1990 and Capan-2 cell lines transfected with miR-301a-3p mimics, inhibitors or negative control. The X-axis indicates the number of hours after transfection. Data represent the mean \pm SEM of the three independent experiments performed in triplicate.

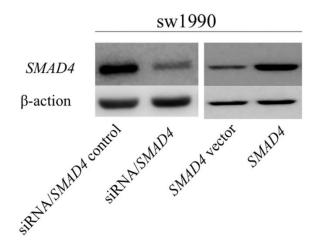


Figure S2: *SMAD4* gene expression was examined via Western blot after *SMAD4* overexpression or knockdown in sw1990 cell.

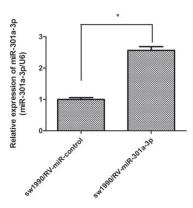


Figure S3: To validate the transfection efficiency of miR-301a-3p in retrovirus-mediated sw1990/ miR-301a-3p and sw1990/ miR-control stable cell lines, qRT-PCR was utilized to measure their miR-301a-3p expression. U6 was used for normalization. Data are shown as mean \pm SEM of three independent experiments. *P<0.05 $_{\circ}$

Table S1: Association between miR-301a-3p ISH score level and PDAC samples in the tissue microarray(n=90)

| Tissue microassay | | miR-301a-3p expression | P value |
|-------------------------|-----------------|------------------------|---------|
| Region | High expression | Low expression | 0.05 |
| Tumor | 46 | 44 | |
| Corresponding non-tumor | 32 | 58 | |

Table S2 Association between SMAD4 IHC score level and PDAC samples in the tissue microarray(n=90)

| Tissue microassay | | SMAD4 expression | P value |
|-------------------------|-----------------|------------------|---------|
| Region | High expression | Low expression | 0.008 |
| Tumor | 24 | 66 | |
| Corresponding non-tumor | 41 | 49 | |